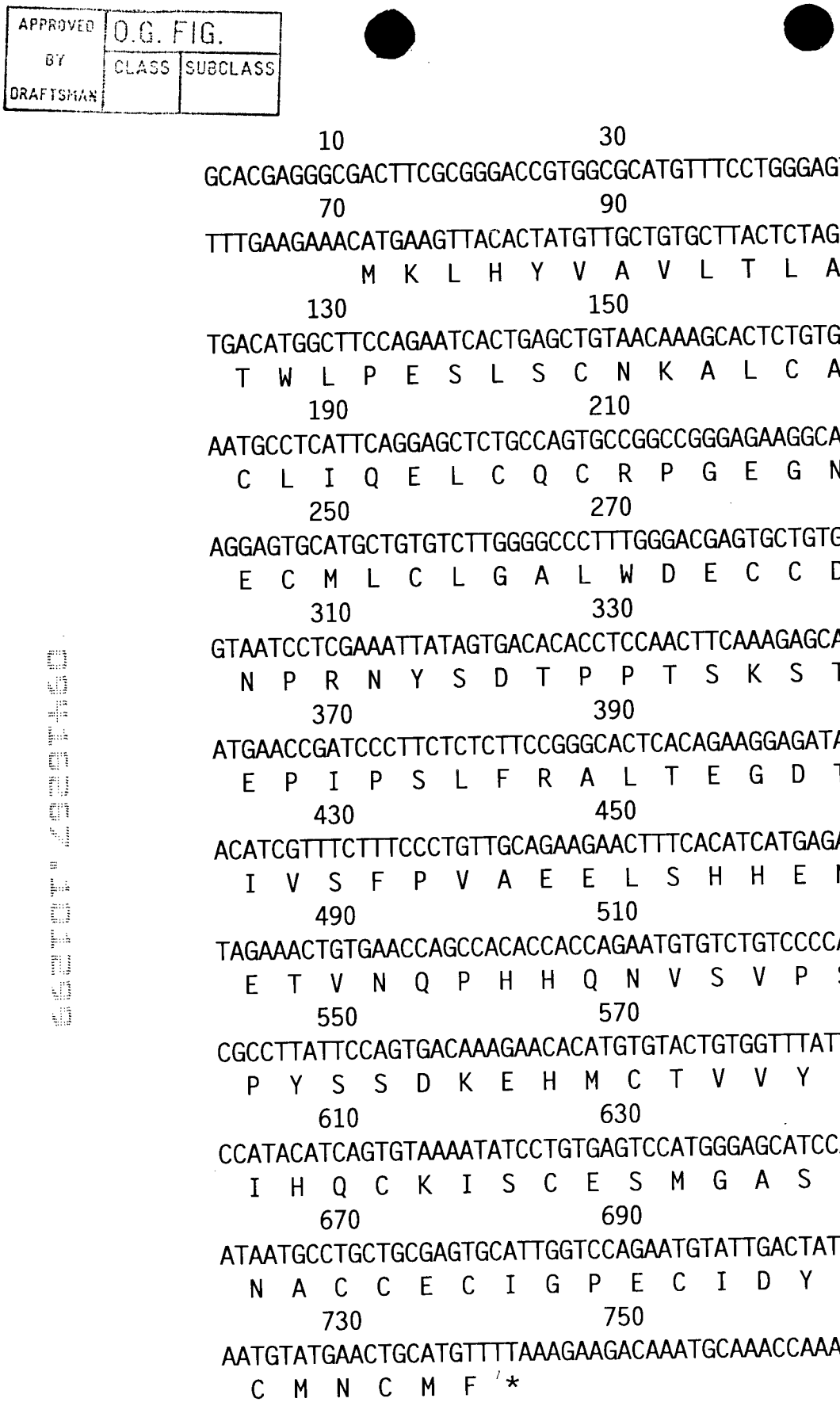

FIG. 1

```
                .         .         .         .         .
  1 MKLHYVAVLTLAILMFLTWLPESLSCNKALCASDVSKCLIQELCQCRPGE  50
    |.| :  |: :. :  :..|::. :||..:|:| |||||| : |||: .:
  1 mqllcyfvilfvglapwsslanddgcnevvcgsvvskclitqscqcklnd  50
                .         .         .         .         .
 51 GNCSCCKECMLCLGALWDECCDCVGMCNPRNYSDTPPTSKSTVEELHEPI 100
      | |||:|: |||.|: |||:|::|| .:.   .. |.:|.:::: |.:
 51 ..chcckdclnclgelyieccgcldmcpkhkdvlpsltprseigdi.egv  97
                .         .         .         .         .
101 PSLFRALTEGDTQLNWNIVSFPVAEELSHHENLVSFLETVNQPHHQNVSV 150
    |.|| .||.:|.: .|..:.|.: .::.:: . ..  :..|
 98 pelfdtltaedde.gwstirfsmragfkqrvqggasgdagn......... 137
                .         .         .         .         .
151 PSNNVHAPYSSDKEHMCTVVYFDDCMSIHQCKISCESMGASKYRWFHNAC 200
      .|.::  :|.   :|||:|.:.|:. :.|:  |||||||.|||||::|
138 ..gngngnagsagvtlctviyvnscirankcrqqcesmgassyrwfhdgc 185
                .         .
201 CECIGPECIDYGSKTVKCMNC                               221
    |||:|.:|::|| .. :|..|
186 cecvgenclnyginesrcrgc                               206
```

FIG.2

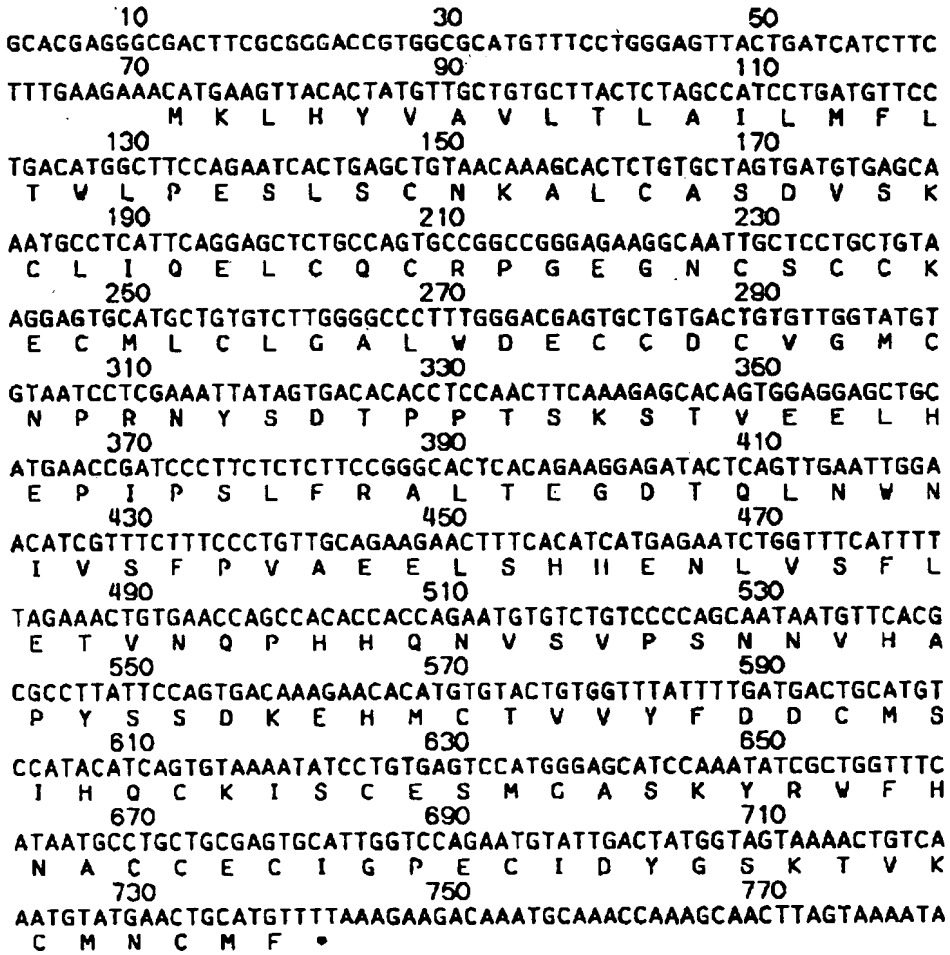
10 30 50
GCACGAGGGCGACTTCGCGGGACCGTGGCGCATGTTTCCTGGGAGTTACTGATCATCTTC
70 90 110
TTTGAAGAAACATGAAGTTACACTATGTTGCTGTGCTTACTCTAGCCATCCTGATGTTCC
M K L H Y V A V L T L A I L M F L
130 150 170
TGACATGGCTTCCAGAATCACTGAGCTGTAACAAAGCACTCTGTGCTAGTGATGTGAGCA
T W L P E S L S C N K A L C A S D V S K
190 210 230
AATGCCTCATTCAGGAGCTCTGCCAGTGCCGGCCGGGAGAAGGCAATTGCTCCTGCTGTA
C L I Q E L C Q C R P G E G N C S C C K
250 270 290
AGGAGTGCATGCTGTGTCTTGGGGCCCTTTGGGACGAGTGCTGTGACTGTGTTGGTATGT
E C M L C L G A L W D E C C D C V G M C
310 330 350
GTAATCCTCGAAATTATAGTGACACACCTCCAACTTCAAAGAGCACAGTGGAGGAGCTGC
N P R N Y S D T P P T S K S T V E E L H
370 390 410
ATGAACCGATCCCTTCTCTCTTCCGGGCACTCACAGAAGGAGATACTCAGTTGAATTGGA
E P I P S L F R A L T E G D T Q L N W N
430 450 470
ACATCGTTTCTTTCCCTGTTGCAGAAGAACTTTCACATCATGAGAATCTGGTTTCATTTT
I V S F P V A E E L S H H E N L V S F L
490 510 530
TAGAAACTGTGAACCAGCCACACCACCAGAATGTGTCTGTCCCCAGCAATAATGTTCACG
E T V N Q P H H Q N V S V P S N N V H A
550 570 590
CGCCTTATTCCAGTGACAAAGAACACATGTGTACTGTGGTTTATTTTGATGACTGCATGT
P Y S S D K E H M C T V V Y F D D C M S
610 630 650
CCATACATCAGTGTAAAATATCCTGTGAGTCCATGGGAGCATCCAAATATCGCTGGTTTC
I H Q C K I S C E S M G A S K Y R W F H
670 690 710
ATAATGCCTGCTGCGAGTGCATTGGTCCAGAATGTATTGACTATGGTAGTAAAACTGTCA
N A C C E C I G P E C I D Y G S K T V K
730 750 770
AATGTATGAACTGCATGTTTTAAAGAAGACAAATGCAAACCAAAGCAACTTAGTAAAATA
C M N C M F •

FIG 1

```
  1 MKLHYVAVLTLAILMFLTWLPESLSCNKALCASDVSKCLIQELCQCRPGE 50
    l.l :  l: :. :  :..l::. :ll..:l:l llllll : lll: .:
  1 mqllcyfvilfvglapwsslanddgcnevvcgsvvskclitqscqcklnd 50

 51 GNCSCCKECMLCLGALWDECCDCVGMCNPRNYSDTPPTSKSTVEELHEPI 100
      l lll:l: lll.l: lll:l::ll .:.   .. l.:l.:::: l.:
 51 ..chcckdclnclgelyleccgcldmcpkhkdvlpsltprseigdi.egv 97

101 PSLFRALTEGDTQLNWNIVSFPVAEELSHHENLVSFLETVNQPHHQNVSV 150
    l.ll .ll.:l.: .l..:.l.: .::.:: . ..  :..l
 98 pelfdtltaedde.gwstirfsmragfkqrvqggasgdagn..... 137

151 PSNNVHAPYSSDKEHMCTVVYFDDCMSIHQCKISCESMGASKYRWFHNAC 200
      .l.::  :l.   :lll:l.:.l:. :.l:  lllllll.llllll:l
138 ..gngngnagsagvtlctvlyvnscirankcrqqcesmgassyrwfhdgc 186

201 CECIGPECIDYGSKTVKCMNC 221
    lll:l.:l::ll .. :l..l
186 cecvgenclnyginesrcrgc 206
```

FIG 2